Figure 1.

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AAGGTGATGGTTGGTCAAAAGTTGGACTCGATGATCTCAGAGTTTTTTTCCAGCCTTAAT     120
AATTCTATGAATTCTGTAATTTTATTCTTGATCTTTTTGAGCGAAGTTTGTTTGGGGATT     180
TTAGTTTGGTTTCCCTGTCACTGTTTTCTTTCCTTGAAACTGACTTTCATTTGCAACATG     240
AGAATTGCTGTATTTGTCAGGTTACAAGTAGTGCAATGGCTGCTTAGAAGTAGTGAGAAA     300
CATTTAGGGAAATACTGGAGTGAAGCAAACACAGTGGTACTGCCAAACTGTAGCTTTGGG     360
ATTTGAGGAGCCACAGAGTTGTATATAAATTTGTTTAATGATATCCTGCCCCTGCCTTCC     420
ATTAATTGCTTGTTTTATGAAACCACTCTTTTTTTTTTTTTTTTTTTTGGCTTCTTCA      480
TATCCTGTGGTAATGAGTTAATGCATTTAGAAGCACATGGCAGAACTAGGAGATCTGTGG     540
ATGACAGTGGTACAGGAGCTCTGAATTTTTTAGATAAACTATGAGAGTGGAAACAGAAAT     600
CTGAGGCTAGTTTCTTGAGCTGACTGTAAATTTTGTGAGAATATTTTCAAGACTACATTA     660
GTTGTGTGTTTGAGGAAAAATAAAATGTTTAAGTTGTCCATTCCTTGAAACCTCCCGACC     720
GGG                                                              723
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Figure 2.

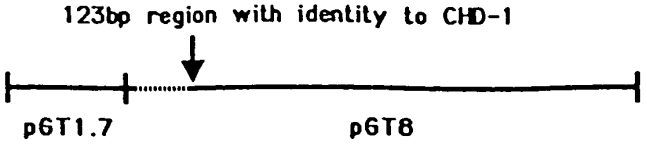

Figure 3.

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C   CHD-W     ATTTTACCTG ATGATCCAGA TAAGAAACCC CAGGCTAAGC AGTTACAGAC
GT  CHD-W     ATTTTACCTG ATGACCCAGA TAAGAAACCA CAGGCAAAGC AGTTGCAGAC

M   CHD-1     CAAAAAACCA CAAGCAAAAC AGTTACAGAC CCGTGCAGAC TACCTCATCA
C   CHD-1A    CAAGAAACCC CAGGCAAAGC AGCTACAGAC CCGTGCAGAC TACCTCATTA
C   CHD-W     CAAGAAACCC CAGGCTAAGC AGTTACAGAC CCGTGCAGAT TACCTCATTA
GT  CHD-W     CAAGAAACCA CAGGCAAAGC AGTTGCAGAC CCGTGCAGAT TACCTCATTA

M   CHD-1     AACTACTTAG CAGAGATCTT GCAAAAAGAG AGGCTCAGAG ACTTTGTGGT GCG
C   CHD-1A    AATTACTGAA TAAAGACCTT GCAAGAAAGG AAGCACAAAG GCTTGCTGGT GCA
C   CHD-W     AATTACTGAA TAAAGACCTT GCAAGAAAGG AAGCACAGAG ACTTGCTGGT GCA
GT  CHD-W     AATTACTGAA TAAAGACCTT GCAAGAAAAG AAGTGCAAAG ACTTACTGGT GCA


M   CHD-1          ILPDDPDKKPQAKQLQTRADYLIKLLSRDLAKREAQRLCGA
C   CHD-1A         ILPDDPDKKPQAKQLQTRADYLIKLLNKDLARKEAQRLAGA
C   CHD-W          ILPDDPDKKPQAKQLQTRADYLIKLLNKDLARKEAQRLAGA
GT  CHD-W          ILPDDPDKKPQAKQLQTRADYLIKLLNKDLARKEVQRLTGA
                   ***************************  **** * *** **
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Figure 4.

08+0220 "29E2V080

Figure 5.

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 151  GGAGTACCAG AAAGGGGATT CTTGACCTAC ACCTTGTAAC CTGAGTGGAC
 201  TTTCTTTTTA ACTTCTTAAT ACTTACAATG AATGGGCACA GTGATGAAGA
 251  AAGTGTAAGA AACAGCAGTG GAGAGTCAAG CAGATCAGAT GATGATTCTG
 301  GGTCAGCTTC AGGTTCTGGA TCTGGTTCAA GCTCTGGAAG CAGTAGCGAT
 351  GGAAGTAGCA GCCAGTCAGG TAGCAGTGAC TCTGAATCTG GTTCAGAGTC
 401  AGGCAGTCAA TCCGAATCAG AGTCTGACAC ATCTAGAGAG AAGAAACAAG
 451  TTCAAGCTAA ACCTCCGAAA GCTGACGGAT CTGAGTTTTG GAAGTCCAGT
 501  CCAAGCATAC TTGCTGTACA GAGATCAGCA GTGCTCAAGA AGCAACAGCA
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3151  AAGAACCTGA AGGAGAAGAA CAGGAGCCCC AGGAAATGGA TATAGATGAA
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3251  AGGGGATGAG TTGCTTTCAC AGTTCAAGGT GGCGAACTTT TCCAATATGG
3301  ATGAAGATGA TATTGAGTTG GAACCAGAAA GAAATTCAAG AAATTGGGAA
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                              **
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Figure 6.

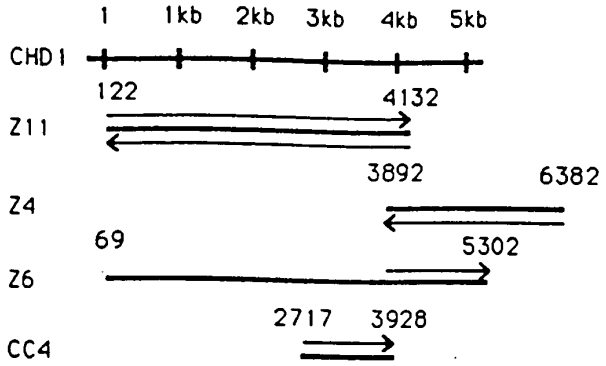

Figure 7.

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CHD-W      1     GATGGGATTGTTTCAGTGAAACATCCACATAAAAAAATAAAAGCAGAAA
                 D  G  I  V  S  V  K  H  P  H  K  K  I  K  A  E

                 K  E  N  E  E  K  P  E  P  D  I  G  I  K  K  E  A
CHD-1A     51    AAAGAAAATGAAGAAAAAGCCTGAGCCAGATATTGGTATAAAGAAGGAAGCT
CHD-W      51    AAAGAAAATGAAGAAAAAGATGAGCCAGAGATTGGTATAAAGAAGGAAGCT
                 K  E  N  E  E  K  D  E  P  E  I  G  I  K  K  E  A

                 E  E  K  R  E  T  K  E  K  E  N  K  R  E  L  K  R
CHD-1A    101    GAAGAAAAAAGAGAGACAAAAGAGAAGGAAAATAAAAGGGAATTGAAAAGG
CHD-W     101    GGAGAAAAAAGAGAGACAAAAGAAAAGGAAAATAAGA
                 G  E  K  R  E  T  K  E  K  E  N  K

                 E  K  K  E  K  E  D  K  K  E  L  K  E  K  D  N  K
CHD-1A    151    GAGAAAAAAGAAAAAGAGGATAAGAAAGAATTAAAAGAAAAAGATAATAAA

                 E  K  R  E  N  K  V  K  E  S  T  Q  K  E  K  E  V
CHD-1A    201    GAAAAGAGAGAAAACAAAGTAAAAGAATCCACACAGAAAGAAAAAGAAGTG

                 K  E  E  K
CHD-1A    251    AAGGAAGAGAAG
```

Figure 8.

```
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        10         20         30         40         50         60         70         80
AGTGATTCAG AGAATGGACA CCACAGGGAA AACTGTACTA CATACAGGCT CTACTCCTTC AAGCTCAACA CCTTTTAATA
        90        100        110        120        130        140        150        160
AGGAAGAGTT ATCAGCAATT TTGAAGTTTG GTGCTGAGGA ACTTTTTAAA GAACCTGAAN NNGAAGAAGA GGAGCCTCAG
       170        180        190        200        210        220        230        240
GAGATGGATA TAGATGAAAT CCTGAAGAGG NCTGAAACTC GAGAAAATGA GTCAGGCCCA TTAACTGTAG GAGATGAGTT
       250        260        270        280        290        300        310        320
ACTTTCACAG TTCAAGGTAG CTAACTTTTC CAATATGGAT GAAGATGACA TTGAATTGGA ACCAGAACAA AATCTAAGAA
       330        340        350        360        370        380        390        400
ACTGGGAAGA AATCATTCCA GAAGTTCAGT GGCGACGAAT AGAGGGGNNG GAAAGACAAA AAGAACTTGA AGAAATATAT
       410        420        430        440        450        460        470        480
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       490        500        510        520        530        540        550        560
TTCTGGATCT GATAGTGATT CCATCTCAGA AAGAAAACGA CCAAAAAAAC GTGGACGACC ACGAACTATT CCCCGTGAAA
       570        580        590        600        610        620        630        640
ACATTAAAGG ATTTAGTGAT GCAGAGATTA GACGATTTAT CAAGAGTTAC AAGAAATTTG GTGGCCCAGT TGAAAGGTTA
       650        660        670        680        690        700        710        720
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       890        900        910        920        930        940        950        960
CCTTCAGATC CAGAAGAAAG GAAAAGATAT GTCATCCCAT ACCACACCAA AGCAGCTCAT TTTGATATAG ATTGGGGTAA
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      1290       1300       1310
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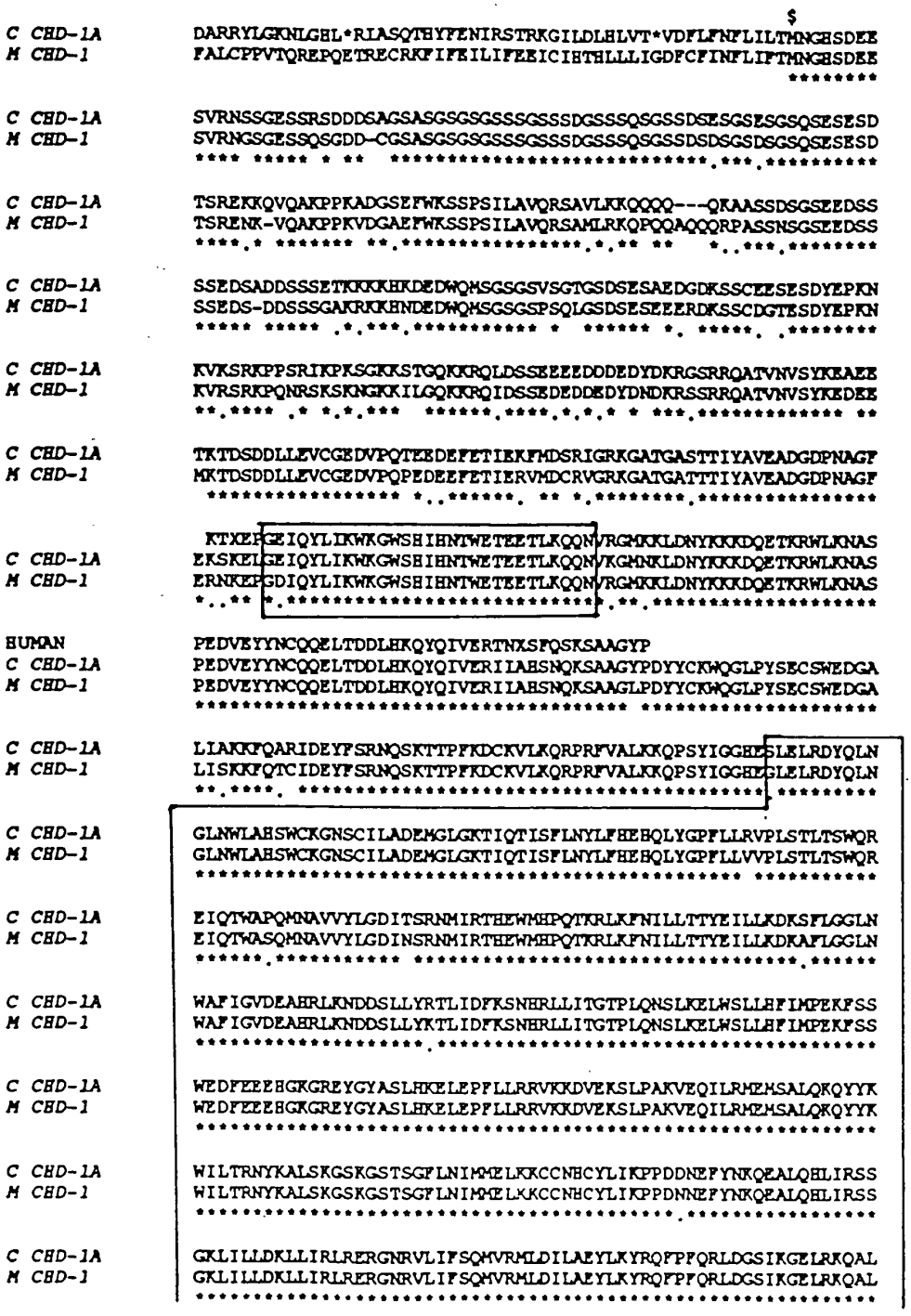
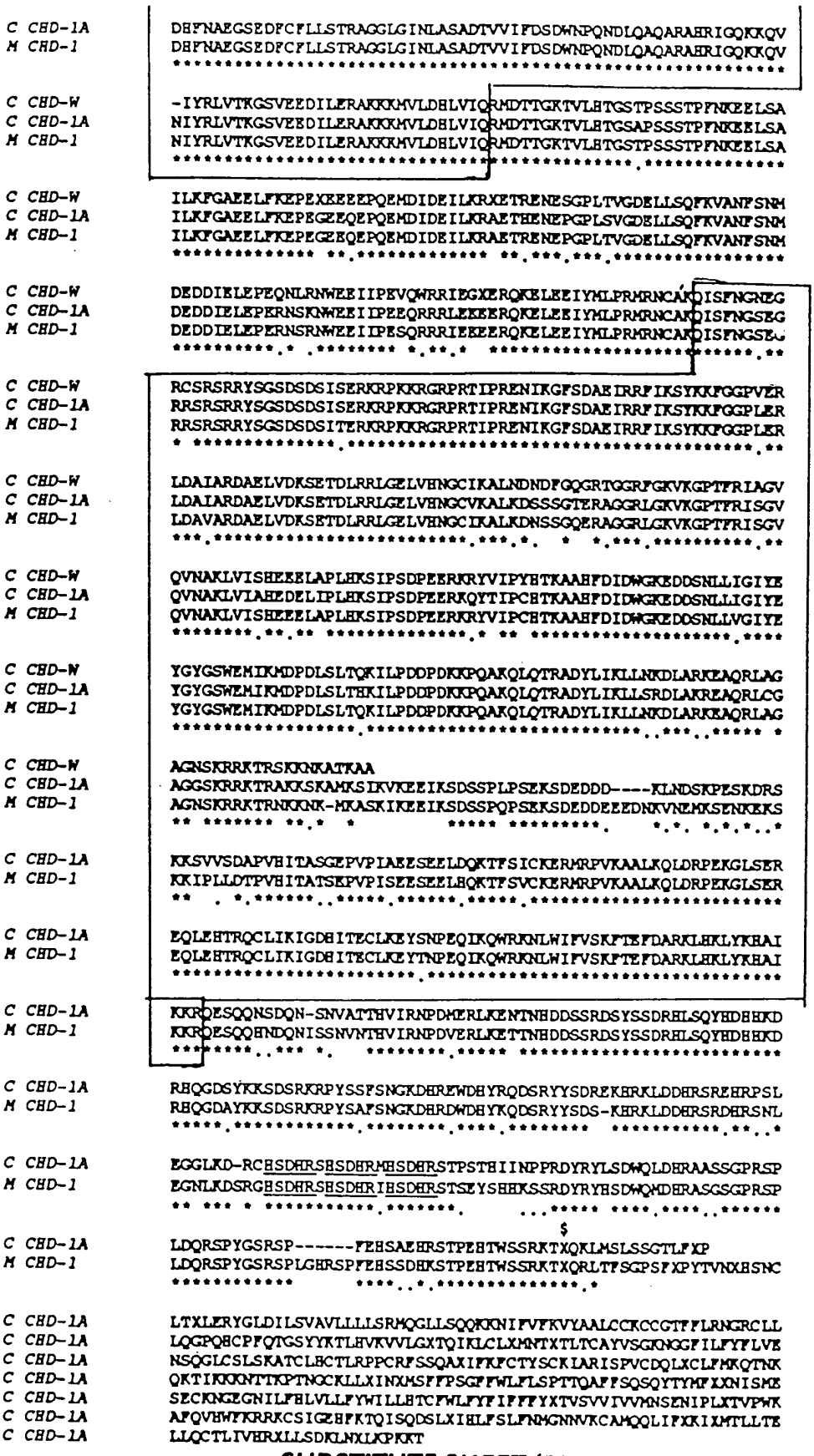
Figure 9.





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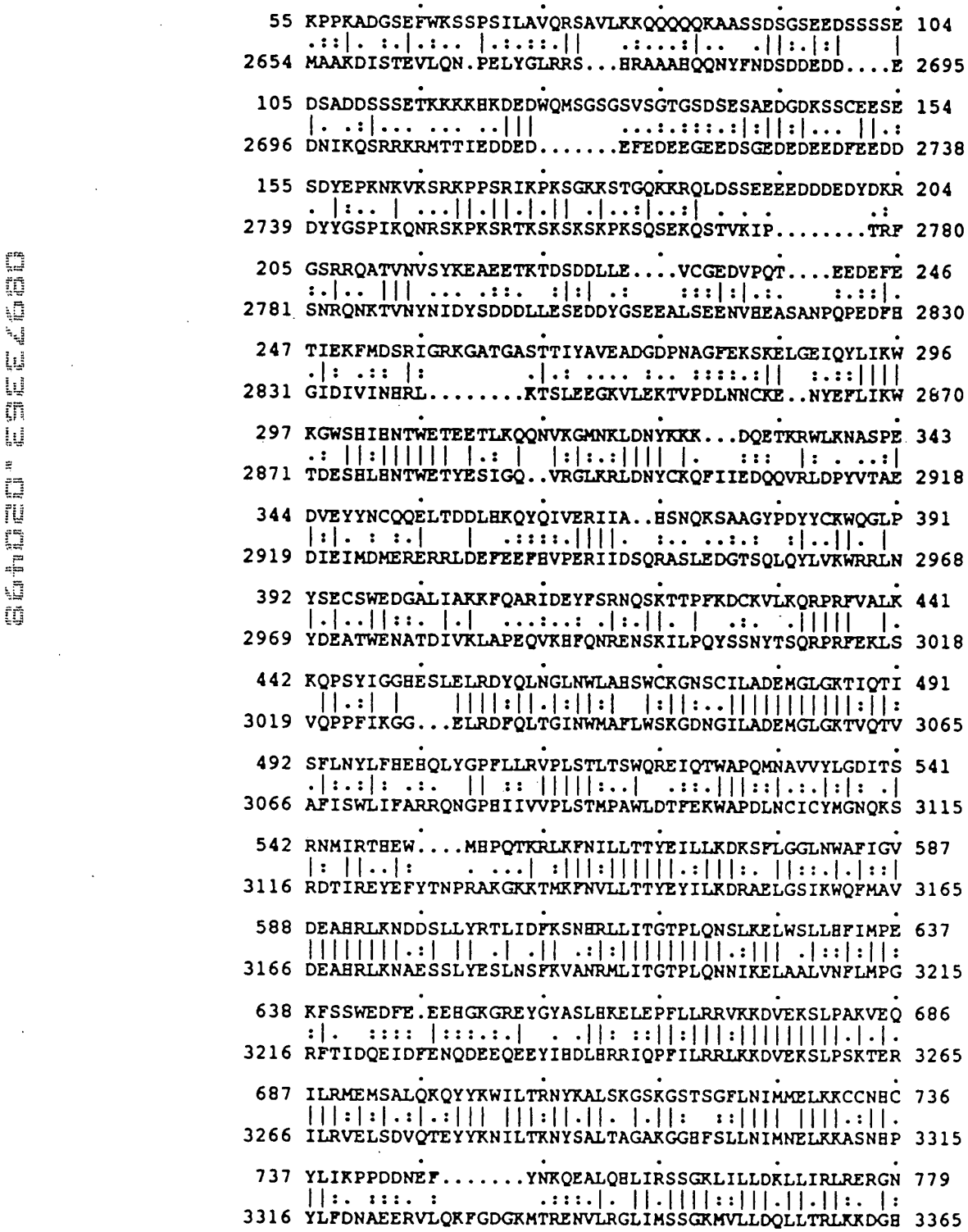

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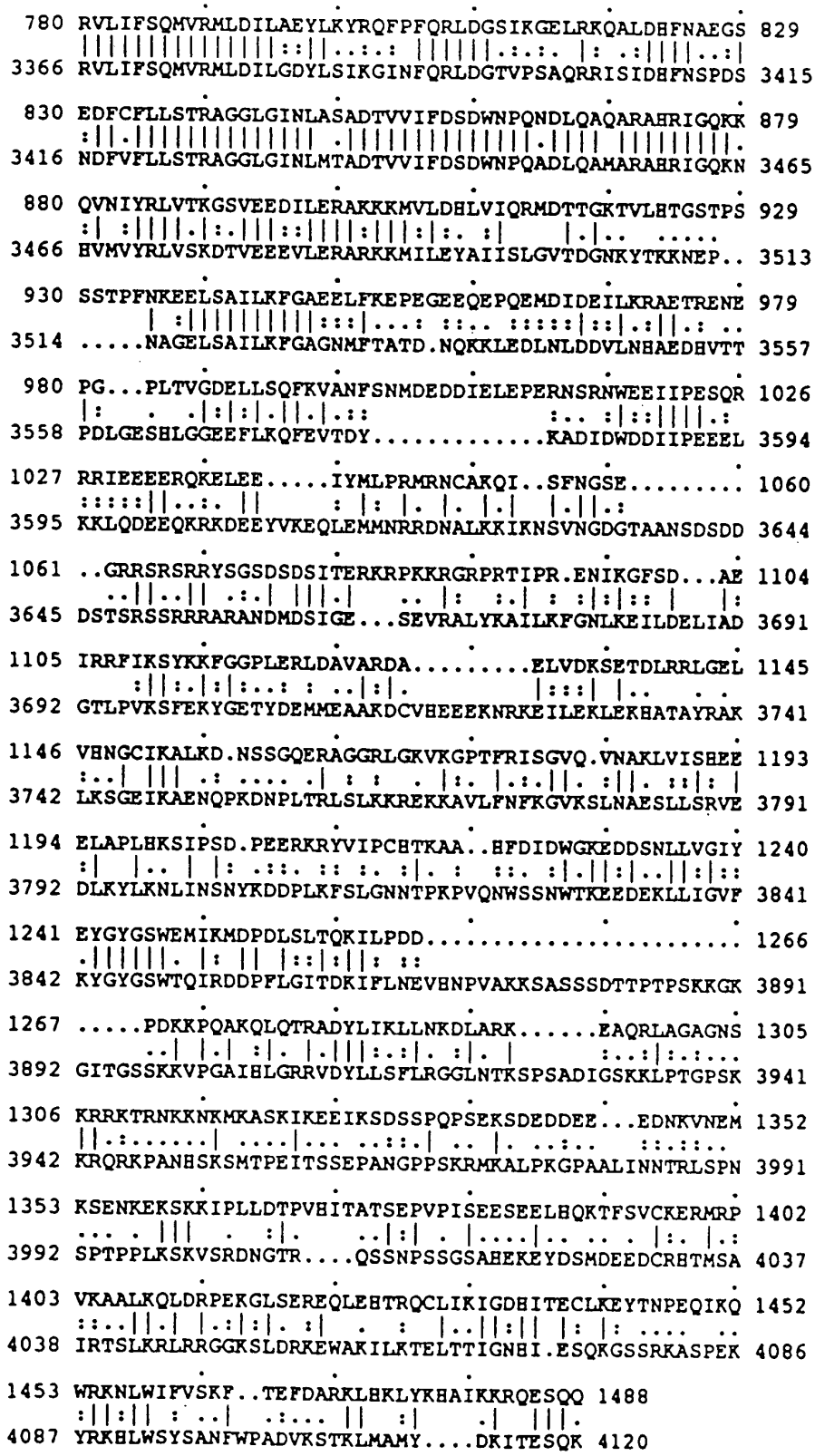

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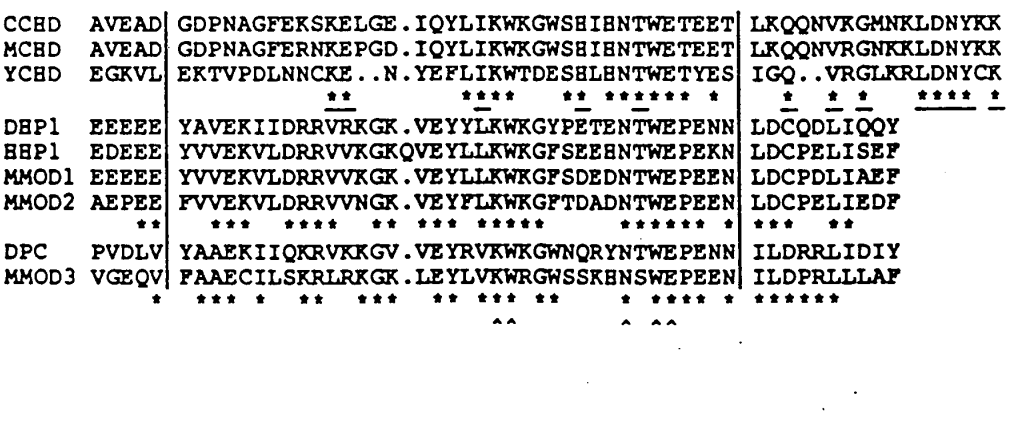

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Figure 12.

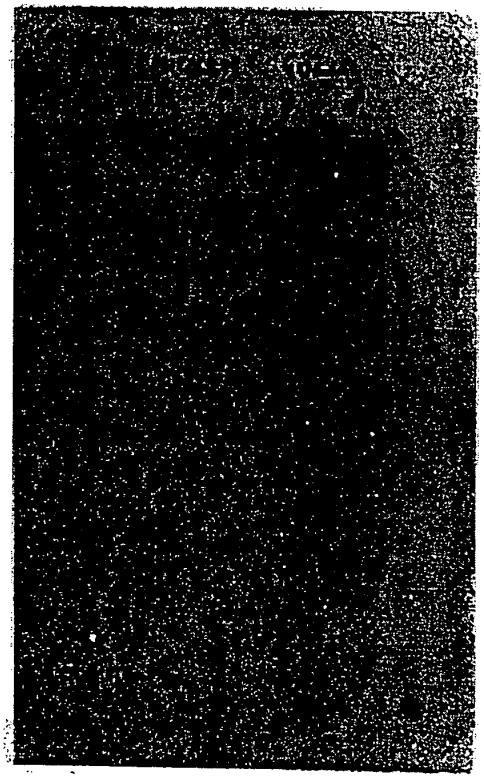

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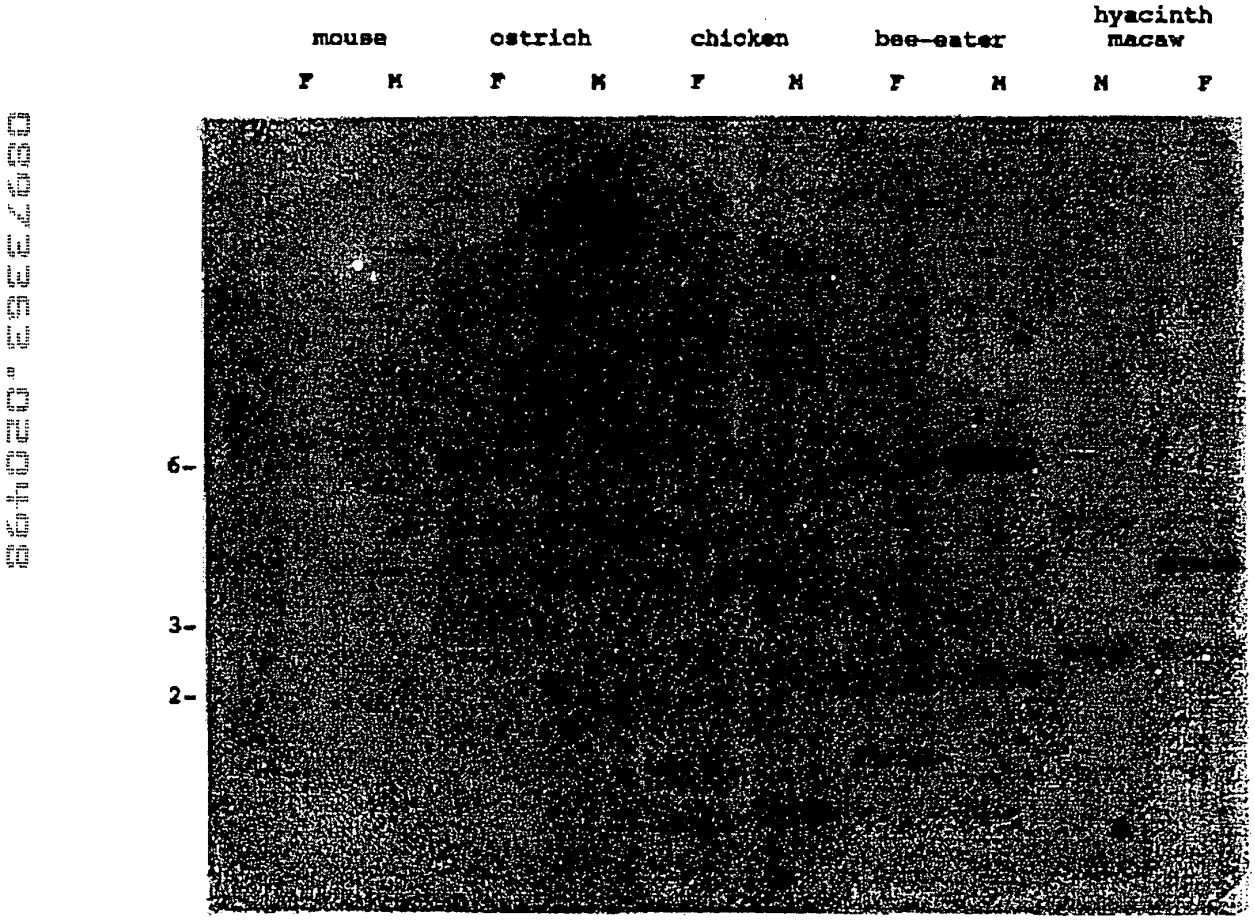
Figure 13.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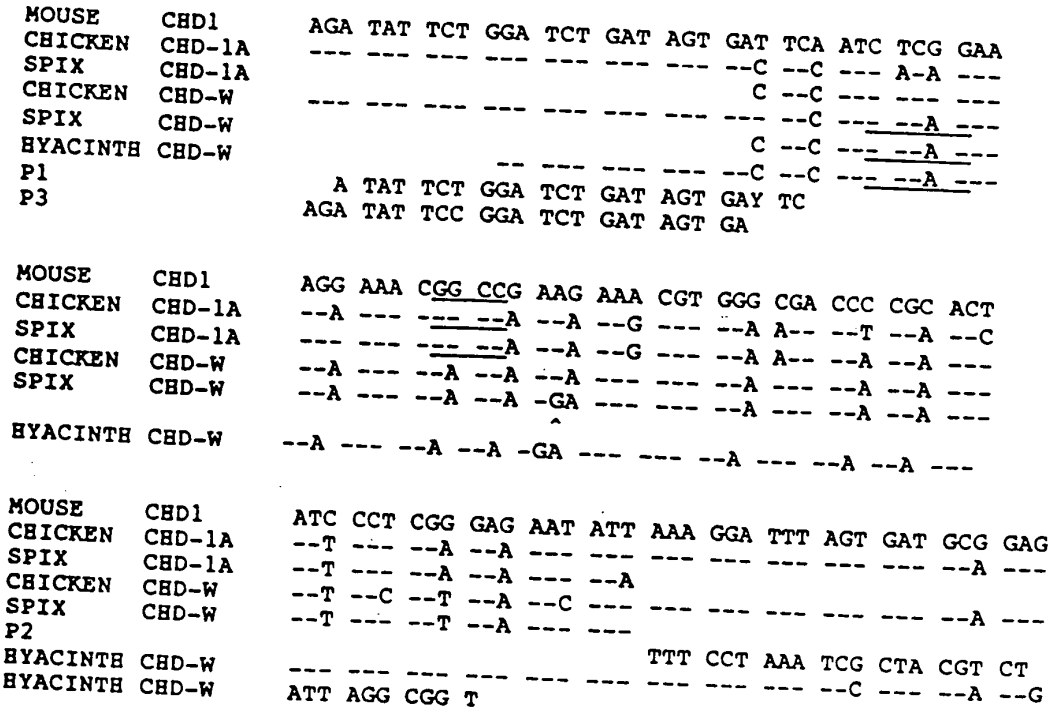
ostrich
chicken
bee-eater
hyacinth macaw
F M F M F M F M M F
6-
3-
2-

Figure 14.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SPIX CHD-1A
CHICKEN CHD-W
SPIX CHD-W
HYACINTH CHD-W
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A TAT TCT GGA TCT GAT AGT GAY TC
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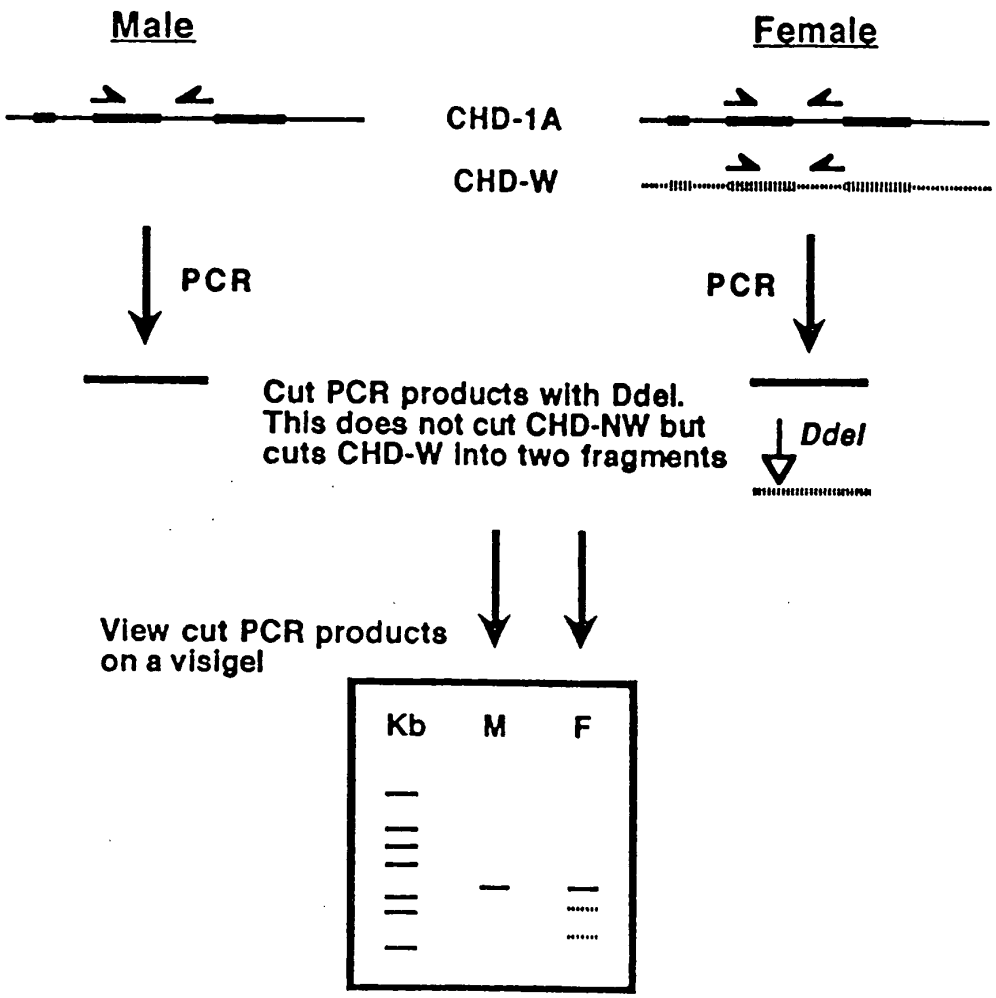
AGG AAA CGG CCG AAG AAA CGT GGG CGA CCC CGC ACT
HYACINTH CHD-W
--A --- --A --A -GA --- --- --A --- --A --A ---
ATC CCT CGG GAG AAT ATT AAA GGA TTT AGT GAT GCG GAG
P2
TTT CCT AAA TCG CTA CGT CT
ATT AGG CGG T

Figure 15.



SUBSTITUTE SHEET (RULE 26)

Figure 16.

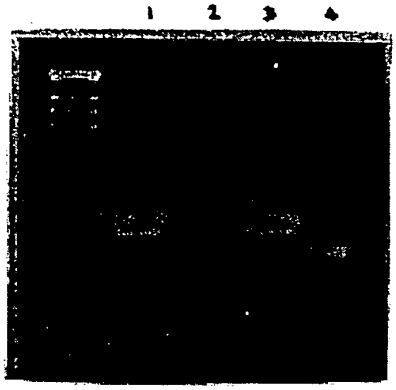
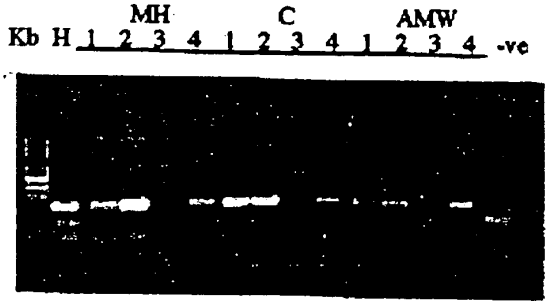



Figure 17.